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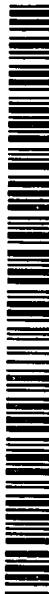
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(54) Title: RECOMBINANT PROTEINS CONTAINING SHIGA-LIKE TOXIN AND VASCULAR ENDOTHELIAL GROWTH FACTOR FRAGMENTS

(57) Abstract: The present invention is directed to an isolated nucleic acid encoding a fusion protein comprising (1) the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and (2) human vascular endothelial growth factor, or a truncated or mutated version thereof; wherein the fusion protein possesses ribosome inactivating activity and ability to bind to cellular VEGF receptors. The present invention is also directed to polypeptides the above combination of toxin and growth factor, as well as expression vectors and transformed cells incorporating the above nucleic acid. The invention is also directed to pharmaceutical compositions and methods for treating patients suffering from diseases relating to angiogenesis.

RECOMBINANT PROTEINS CONTAINING SHIGA-LIKE TOXIN AND VASCULAR ENDOTHELIAL GROWTH FACTOR FRAGMENTS

5

BACKGROUND OF THE INVENTION

1. Field of the Invention

The present invention relates to recombinant nucleic acid molecules and recombinant fusion proteins, and more particularly to Shiga-like toxin-vascular endothelial growth factor fusion proteins and recombinant DNA molecules coding for such fusion proteins. The present invention also relates to bacterial vectors containing the above recombinant nucleic acid molecules, methods of producing the above fusion proteins, and their use in therapeutic treatments.

2. Description of the Related Art

Angiogenesis is a tightly controlled process of growing new blood vessels (see, Folkman & Shing, 1992; Hanahan, 1997, for reviews). Under normal circumstances angiogenesis occurs only during embryonic development, wound healing and development of the corpus luteum. However, angiogenesis occurs in a large number of pathologies, such as solid tumor and metastasis growth, various eye diseases, chronic inflammatory states, and ischemic injuries (see, Folkman, 1995, for review). Thus, growing endothelial cells present unique targets for treatment of several major pathologies.

The crucial positive regulator of angiogenesis is vascular endothelial growth factor (VEGF) also known as vascular permeability factor (see, Neufeld, et al, 1999 for reviews). VEGF is a secreted dimeric glycoprotein that, as a result of alternative splicing, may consist of polypeptides with 121, 145, 165, 189 and 206 amino acid residues. VEGF is expressed by normal and tumor cells and the control of VEGF expression appears to be

regulated on several levels (see, Claffey & Robinson, 1996, Veikkola & Alitalo, 2000, for reviews). Expression of VEGF is upregulated in response to hypoxia and nutritional deprivation suggesting a feedback loop between tumor and metastasis growth and the ability of tumor cells to induce host angiogenic responses.

5 The action of VEGF on endothelial cells is mediated by tyrosine kinase flt-1 and KDR/flk-1 receptors, also known as VEGFR-1 and VEGFR-2 (see, Terman, & Dougher-Vermazen, 1996; Veikkola, et al, 2000, for review). These receptors are preferentially expressed on endothelial cells. There are reports that endothelial cells at the sites of angiogenesis express significantly higher numbers of KDR/Flk1 receptors than quiescent
10 endothelial cells (Brown, et al., 1993, 1995; Plate, et al., 1993; Detmar, et al., 1994; Couffinhal, et al., 1997). The receptors are single span transmembrane protein tyrosine kinase that belong to the immunoglobulin superfamily and contains seven Ig-like loops in the extracellular domain and shares homology with the receptor for platelet-derived growth factor. VEGF binding to these receptors induces receptor dimerization followed by
15 tyrosine phosphorylation of the SH2 and SH3 domains in the dimer (see, Neufeld, et al., 1994 for review). KDR/Flk1-VEGF complex is internalized via receptor-mediated endocytosis (Bikfalvi, et al., 1991).

Several groups reported that targeting of either VEGF or KDR/flk-1 inhibits angiogenesis and angiogenesis-dependent processes (Kim, et al., 1993; Millauer, et al.,
20 1994; Saleh, et al., 1996; Aiello, et al., 1995). On the other hand, direct injection of VEGF or a plasmid encoding VEGF into ischemic tissues in a model system promoted development of microvasculature and improved recovery after ischemic injury or balloon angioplasty (Asahara, et al., 1996). Taken together, these results leave little doubt that VEGF and KDR/Flt1 play crucial roles in angiogenesis. Although these experiments provided a
25 "proof-of-principle" that VEGF-toxin conjugates or fusion proteins may work *in vivo*,

further development of DT-VEGF constructs is doubtful, because of the renal and liver toxicity of DT-containing fusion proteins (see, for example, Vallera et al., 1997).

Since VEGF binds specifically to endothelial cells, this growth factor provides a unique opportunity for targeted drug delivery to the sites of angiogenesis. It was

5 demonstrated that catalytically active forms of diphtheria toxin covalently linked or fused via recombinant DNA technology to recombinant VEGF165 and/or VEGF121 are selectively toxic against cells expressing KDR/flk-1 receptors and also suppressed angiogenesis in vivo (Ramakrishnan, et al., 1996; Olson et al., 1997; Arora, et al., 1999).

It is advantageous to use VEGF for targeting toxins that are "natural killers" of
10 endothelial cells. Shiga-like toxin 1 produced by E. coli O157:H7 is such a "natural killer" for endothelial cells. Damage to endothelial cells caused by Shiga-like toxins 1 plays a causative role in the pathogenesis of hemorrhagic colitis (HC) and hemolytic uremic syndrome (HUS) induced by E. coli O157:H7 (Obrig, et al., 1987, 1993; Richardson, et al., 1988; Kaplan, et al., 1990).

15 Shiga-like toxin 1 (SLT-1) is composed of a single copy of a 32 kDa A-subunit associated with a ring shaped pentamer of receptor-binding 7 kDa B-subunits. B-subunits bind SLTs to the cellular receptor globotriaosylceramide known as Gb₃ (Obrig et al., 1993). This receptor is found on many cell types including endothelial cells (Obrig et al., 1993). After binding to the cell surface receptor, SLT is endocytosed and A-subunit is
20 cleaved into A₁ (27.5 kDa) and A₂ (4.5 kDa) forms that are linked by disulphide bond (Olsnes et al., 1981). Processed A subunit is a specific N-glycosidase that inactivates ribosomes by cleaving off a single adenine residue in the position 4324 from 5' terminus of 28S rRNA of 60S ribosome subunit (Saxena et al., 1989). The cleavage of A₄₃₂₄ from 28S rRNA inactivates ribosomes by inhibiting binding of the elongation factor (EF-25 1)/aminocyl-tRNA complex to ribosomes, resulting in the inhibition of the protein

synthesis. As with other ribosome-inactivating agents, the subsequent cytostatic and cytotoxic effects might arise as a cellular response to inactivation of a relatively small proportion of ribosomes through ribotoxic stress response (Iordanov et al., 1997).

Alternatively, cytostatic and cytotoxic effects might arise as a cellular response to a

- 5 massive collapse of protein synthesis due to inactivation of a large number of ribosomes.
- It is important that the unprocessed, full length A subunit as well as various truncated A subunits retain significant N-glycosidase activity (Haddad, et al., 1993; Al-Jaufy, et al., 1994, 1995). Furthermore, fusion proteins containing unprocessed, full length A subunit as well as various truncated A subunits fused to N-terminus of CD4 retain N-glycosidase
- 10 activity and are cytotoxic for cells expressing HIV-1 gp120-gp41 complex (Al-Jaufy, et al., 1994, 1995).

Since Shiga-like toxin is a "natural" killer of endothelial cells it is advantageous to

deliver enzymatically active full-length, truncated or mutated A subunit into endothelial cells in order to inhibit their growth and/or kill them. To avoid damage to other cell type

- 15 the enzymatically active full-length, truncated or mutated A subunit should be delivered into target cells by endothelial cell specific growth factor such as VEGF. Therefore, it is an object herein to provide effective recombinant DNA methods for the production of fusion proteins containing enzymatically active full-length, truncated or mutated A subunit fused to full-length, truncated or mutated VEGF that retain ability to bind to VEGF
- 20 receptors.

SUMMARY OF THE INVENTION

In one aspect, the present invention is directed to an isolated nucleic acid encoding a fusion protein comprising: (1) the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and (2) human vascular endothelial growth factor, or a

truncated or mutated version thereof; wherein the fusion protein possesses ribosome inactivating activity.

In another aspect, the present invention is directed to an isolated polypeptide comprising: (1) the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and (2) human vascular endothelial growth factor, or a truncated or mutated version thereof; wherein the isolated polypeptide possesses ribosome inactivating activity.

In another aspect, the present invention is directed to an expression vector, comprising: (1) a nucleic acid encoding a fusion protein comprising the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and human vascular endothelial growth factor, or a truncated or mutated version thereof; and (2) a promoter sequence operably linked to the nucleic acid to allow expression of the nucleic acid.

In another aspect, the present invention is directed to a bacterial cell transformed with the above expression vector.

In yet another aspect, the present invention is directed to a method of inactivating ribosomes in a cell, comprising the steps of: (a) contacting a cell with a polypeptide comprising: (1) the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and (2) human vascular endothelial growth factor, or a truncated or mutated version thereof; under conditions which permit the polypeptide to be internalized into the cell and inactivate ribosomes in the cell.

In yet another aspect, the present invention is directed to a composition for inhibiting endothelial cell growth in a patient, comprising: (A) a fusion protein comprising the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and human vascular endothelial growth factor, or a truncated or mutated version thereof, the

fusion protein possessing ribosome inactivating activity; and (B) a pharmaceutically acceptable carrier.

In yet another aspect, the present invention is directed to a method of treating a patient suffering from a pathophysiological condition that depends on angiogenesis,

5 comprising: providing to the patient an effective amount of a composition comprising a fusion protein comprising the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and human vascular endothelial growth factor, or a truncated or mutated version thereof, the fusion protein possessing ribosome inactivating activity; and a pharmaceutically acceptable carrier.

10 In addition, the proteins and pharmaceutical compositions of the present invention may be used either alone, or in combination with other treatments for diseases related to angiogenesis, particularly treatments whose efficacy is enhanced by decrease in oxygen or nutrient supplies that would arise from damage to endothelium caused by said protein and pharmaceutical compositions.

15 These and other aspects will be described in more details in the following detailed description of the invention.

BRIEF DESCRIPTION OF THE FIGURES

The invention will be more fully understood from the following detailed description taken in conjunction with the accompanying figures in which:

Figure 1 is a schematic representation of SLT-VEGF/L, catalytically inactive SLT-VEGF/Lci, and SLT-VEGF/S proteins. Catalytically inactive SLT-VEGF/Lci was constructed in order to separate the effects of ribosome inactivation from other effects that might be induced by recombinant SLT-VEGF proteins. This protein contains a double mutant A-subunit with Y114S and R170L amino acid substitutions that independently

significantly decrease the enzymatic activity of SLT-1 A-subunit, while not affecting its folding as judged by unchanged antigenic properties (Deresiewicz et al., 1993; Cao et al., 1994). His- and S-tag are used for purification and quantitation. Cleavage site for intracellular protease furin that cleaves A subunits into disulphide bond linked A1-A2 dimers is indicated. Recombinant VEGF121 protein used in control experiments also contains His- and S- tags.

Figure 2 illustrates expression of SLT-VEGF/L, and SLT-VEGF/S proteins in BL21(DE3)pLysS and Origami(DE3)pLysS E. coli strains (termed BL21 and Origami, respectively) and their accumulation in inclusion bodies isolated from respective hosts (Fig. 2, panels A and B). Fig. 2 also illustrates the quality of final preparations of VEGF121 (lane V), SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S proteins obtained after purification from Origami(DE3)pLysS E. coli strain (Fig. 2, panel C). Expression of SLT-VEGF fusion proteins was induced by addition of isopropyl- β -D-thio-galactopyronoside (IPTG). BL21(DE3)pLysS cells were harvested after 3.5 hours IPTG induction for SLT-VEGF/L and after 2 hours of IPTG induction for SLT-VEGF/S at 37°C. Origami(DE3)pLysS cells were harvested after 4 hours IPTG induction for both proteins at 30°C. Soluble fractions (S), inclusion bodies (I), and refolded proteins purified from inclusion bodies were analyzed by SDS-PAGE on 15% gels. Molecular weights of markers in lane M are indicated in kDa.

Figure 3 illustrates that SLT-VEGF/L and SLT-VEGF/S proteins, but not catalytically inactive SLT-VEGF/Lci, inhibit protein synthesis in cell-free translation system. SLT-VEGF/L and SLT-VEGF/S fusion proteins inhibit translation of firefly luciferase mRNA by 99.99% and 99%, respectively at concentration of 100 nM (Fig. 3, panel A). Recombinant VEGF121 protein isolated by the same procedure as SLT-VEGF fusion proteins inhibits translation only ~50% at concentration as high as 1,000 nM (Fig.

3, panel A). SLT-VEGF/L and SLT-VEGF/S inhibited protein synthesis in a dose-dependent manner with 90% inhibition at concentrations 0.04 nM and 2 nM, respectively, while SLT-VEGF/Lci did not inhibit protein synthesis (Fig. 3, panel B). Detected luciferase activities in percents of the VEGF121 control are indicated.

5 Figure 4 illustrates that SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S proteins induce tyrosine phosphorylation of KDR/flk-1 receptors for VEGF in cells overexpressing KDR/flk-1 receptors (293/KDR cells). Tyrosine phosphorylation of KDR/flk-1 receptors was detected by Western blot analysis of the lysates of 293/KDR treated with SLT-VEGF/L, SLT-VEGF/Lci, SLT-VEGF/S, and VEGF121, using anti-phosphotyrosine 10 antibody.

Figure 5 illustrates that SLT-VEGF/L and SLT-VEGF/S proteins target growing PAE/KDR cells that overexpress KDR/flk-1 receptors (open circles) but do not affect control PAE/V cells that do not express KDR/flk-1 receptors (filled circles). PAE/KDR cells and control PAE/V cells lacking KDR/flk-1 receptors were plated at ~5,000 15 cells/well and treated for 72 hours with SLT-VEGF/L (Fig. 5, panel A) or SLT-VEGF/S (Fig. 5, panel B) isolated from Origami(DE3)lysS host. As shown in Fig. 5, SLT-VEGF proteins strongly inhibit growth of PAE/KDR cells overexpressing KDR/flk-1 receptors. This effect is thought to be due to the ribosome-inactivating activity of SLT moiety, because catalytically inactive SLT-VEGF/Lci protein does not affect growth of PAE/KDR 20 and PAE/V cells (Fig. 5, panel C).

Figure 6 illustrates that SLT-VEGF/L fusion protein rapidly activates apoptosis in PAE/KDR cells as judged by DNA degradation (Fig. 6, panel A) and cleavage of α -fodrin (Fig. 6, panel B).

Figure 7 illustrates that SLT-VEGF/L proteins do not target endothelial cells with a 25 low number of KDR/flk-1 receptors (Fig. 7, panel A) and quiescent PAE/KDR cells (Fig.

7, panel B). As estimated by Western blot analysis, human umbilical vein endothelial (HUVE) cells express 30,000-50,000 KDR/flk-1 receptors per cell, and PAE/KDR_{low} cells express ~5,000 KDR/flk-1 receptors per cell. MS1 cells expressed ~20,000 VEGFR-2/cells. HUVE, PAE/KDR_{low}, and MS1 cells were plated onto 24-well plates at densities 5 of 5-10x10³ cells/well and exposed to 2.5 nM SLT-VEGF/L 20 hr later and counted after 72 hrs. Confluent PAE/KDR were maintained at confluence for 3 days, then treated with 20 nM SLT-VEGF/L for 72 hrs. Growing PAE/KDR were exposed to 20 nM SLT-VEGF/L for 5 min; then shifted to fresh culture medium and counted after 72 hrs.

10

DETAILED DESCRIPTION OF THE INVENTION

An object of the present invention is to provide nucleic acid sequences, such as DNA or RNA sequences that code for fusion proteins herein named SLT-VEGF. The SLT-VEGF fusion proteins include a full-length, truncated, or mutated A subunit of Shiga-like bacterial toxin which confers ribosome inactivating activity, and a vascular 15 endothelial growth factor (VEGF) that binds to VEGF receptors. The three nucleic acid sequences and resulting protein sequences are preferably separated by a spacer sequence.

The nucleic acid sequences for the Shiga-like toxin and the VEGF are individually known in the art. However, the inventors have surprisingly found that a combination of these two sequences provides for production of a fusion protein with a unique combination 20 of characteristics. The fusion protein is capable of binding to specific VEGF cellular receptor by virtue of the VEGF domain. The fusion protein is also capable of inactivating ribosomes and inducing apoptosis in endothelial cells overexpressing KDR/flk-1 receptors by virtue of the Shiga-like toxin domain. In combination, these two protein domains provide an effective and highly targeted treatment for diseases relating to angiogenesis.

Still another object of the invention is to provide a pharmaceutical composition for use in inhibition of endothelial cell growth, and containing the SLT-VEGF fusion proteins and a pharmaceutically acceptable carrier. Useful carriers include water, buffered saline, or other pharmaceutically acceptable carrier known in the art. The SLT-VEGF fusion

5 proteins of the pharmaceutical composition are potent cytotoxic or cytostatic agents and are useful in treating of a variety of pathophysiological conditions that depend on angiogenesis, such as solid tumor and metastasis growth, various eye diseases, chronic inflammatory states, and ischemic injuries. In addition, the proteins and pharmaceutical compositions of the present invention may be used either alone, or in combination with

10 other known treatments for diseases related to angiogenesis, particularly treatments whose efficacy is enhanced by decrease in oxygen or nutrient supplies that would arise from damage to endothelium caused by said protein and pharmaceutical compositions.

Yet another object of the invention is to provide recombinant expression vectors harboring the new DNA sequences and transformed bacterial cells containing such

15 recombinant expression vectors. The nucleic acid sequences coding for the fusion protein SLT-VEGF may be inserted into known vectors, such as a bacterial plasmid or viral vector, using materials and methods well known in the art. The nucleic acid construct coding for the SLT-VEGF fusion proteins is inserted into a plasmid such that nucleic acid construct is operatively linked to an inducible promoter sequence, a sequence that encodes

20 tags that simplify purification and quantitation of the fusion protein, and terminator functionality in the selected host. The plasmid is also preferably introduced into a host cell, such as a bacterial cell, in which the promoter is inducibly regulated.

Another object of the invention are methods for inhibiting growth of endothelial cells, and treating a patient suffering from a pathophysiological condition that depends on

angiogenesis, such as solid tumor and metastasis growth, various eye diseases, chronic inflammatory states, and ischemic injuries.

In the fusion protein expressed by the recombinant nucleic acid sequence according to the present invention, the VEGF thereof is suitably selected from full-length or mutants of VEGF121, VEGF165, VEGF189 and VEGF209 capable of binding to high affinity receptors of VEGF. According to a particularly preferred embodiment of the invention, the VEGF is constituted by VEGF121 or truncated VEGF mutants thereof.

As used herein, Shiga-like toxin A subunit (abbreviated herein as SLT) refers to polypeptide having amino acid sequences found in E. coli O157:H7, as well as modified sequences, having amino acid substitutions, deletions, insertions or additions, which still express substantial ribosome inactivating activity. For some applications, such as various control experiments, it may be beneficial to produce SLT lacking ribosome inactivating activity. In particular, such modified SLTs may be produced by modifying the DNA disclosed herein by altering one or more amino acids or deleting or inserting one or more amino acids that may render it more suitable to achieve desired properties of SLT-VEGF fusion proteins. Such properties include but not limited to yield of recombinant protein in bacterial host, ability to bind to cellular VEGF receptor, ability to be internalized via receptor-mediated uptake, intracellular protein synthesis inhibitory activity, overall cytotoxic or cytostatic effects, pharmacokinetics and pharmacodynamics, and stability under various storage and use conditions. Any such protein, or version thereof, that, when fused to VEGF as described herein, that exhibits ribosome inactivating activity and ability to bind to cellular VEGF receptors in standard in vitro or in vivo assays is contemplated for use herein.

As used herein, SLT-VEGF proteins are fusion proteins containing an SLT polypeptide and vascular endothelial growth factor (VEGF), that is reactive with VEGF cell surface receptor.

The resulting SLT-VEGF fusion proteins are useful as cytotoxic or cytostatic agents that target and inhibit growth of endothelial cells and thereby are useful for treating angiogenesis-dependent diseases, including, but not limited to, solid tumor and metastasis growth, various eye diseases, chronic inflammatory states, and ischemic injuries.

As used herein, to target SLT-VEGF protein means to direct it to a cell that expresses VEGF receptors. Upon binding to the receptor SLT-VEGF protein is internalized by the cell and is cytotoxic or cytostatic to the cell.

As used herein, the term active, or reference to the activity of SLT-VEGF proteins or cytotoxic and cytostatic effects of SLT-VEGF proteins, refers to the ability of such proteins to inactivate ribosomes either in vivo or in vitro or respectively, to kill cells or to inhibit cell growth upon VEGF-receptor mediated internalization of SLT-VEGF proteins by the cells. Such activity may be assayed by any method known to those of skill in the art including, but not limited to, the assays that measure protein synthesis, receptor binding, autophosphorylation and internalization and assays that assess cytotoxic and cytostatic effects by measuring the effect of a test compound on cell proliferation, apoptosis and on protein synthesis.

As used herein, VEGF refers to polypeptides having amino acid sequences of native VEGF proteins, as well as modified sequences, having amino acid substitutions, deletions, insertions or additions of the native protein but retaining the ability to bind to VEGF receptors and to be internalized. Such polypeptides include, but are not limited to, VEGF121, VEGF165, VEGF189, VEGF209.

It is understood that differences in amino acid sequences can occur among VEGFs of different species as well as among VEGFs from individual organisms or species.

Reference to VEGFs is also intended to encompass proteins isolated from natural sources as well as those made synthetically, as by recombinant means or possibly by chemical

5 synthesis. VEGF also encompasses mutants of VEGF that possess the ability to target SLT to VEGF-receptor expressing cells and created in order to, for example, retain or increase the activity or stability of the growth factor, to reduce or eliminate disulfide scrambling, or to alter reactivity with various modifying groups (e. g. polyethylene glycol).

10 As used herein, the term "VEGF receptor" is used to refer to receptors that specifically interact with VEGF and transport it into the cell. Included, but not limited to, among these are KDR/flk-1 (VEGF-R1), flt-1 (VEGF-R2).

As used herein, the term "polypeptide reactive with the VEGF receptor" refers to any polypeptide that specifically interacts with VEGF receptor, preferably the high-

15 affinity VEGF receptor, and is transported into the cell by virtue of its interaction with the VEGF receptor.

Unless defined otherwise, all additional technical and scientific terms used herein have the same meaning as is commonly understood by one of skill in the art to which the subject matter herein belongs.

20 Although the invention is by no means limited hereto it will be exemplified in the following mainly with reference to the full length A subunit of Shiga-like bacterial toxin (SLT/L), truncated version thereof (SLT/S), or a catalytically inactive double mutant version thereof (SLT/Lci). Accordingly, the invention will be described in relation to the construction of genetic fusion proteins between SLT/L, or SLT/S, or SLT/Lci and VEGF
25 molecule which will target the fusion protein to specific VEGF receptors, and it will be

demonstrated herein that only fusion proteins containing SLT/L or SLT/S, but not SLT/Lci, exhibit cytotoxic and/or cytostatic effects on endothelial cells.

Fusion proteins denoted SLT-VEGF/L and SLT-VEGF/S consisting of SLT/L or SLT/S linked to VEGF121 inhibit growth of porcine endothelial cells PAE/KDR cells

5 overexpressing KDR/flk-1 receptor for VEGF in a dose-dependent manner with IC₅₀ of ~0.15 nM. At the low nanomolar concentration range SLT-VEGF/L proteins are cytotoxic, killing virtually all PAE/KDR cells after exposure to concentration as low as 2.5 nM. In contrast, SLT-VEGF/S proteins at the low nanomolar concentrations are mostly cytostatic. These effects depend on catalytic activity of SLT moiety in fusion protein that inactivates

10 ribosomes, because catalytically inactive VEGF-SLT/Lci does not affect PAE/KDR cell growth. These effects depend on expression of KDR/flk-1 receptors, because SLT-VEGF/L and SLT-VEGF/S do not affect growth of porcine endothelial cells PAE/V cells that do not express KDR/flk-1 receptors but transfected by control vector. Importantly, SLT-VEGF/L proteins do not affect endothelial cells that express low numbers of

15 KDR/flk-1 receptors or quiescent PAE/KDR cells even at concentration as high as 20 nM. The results demonstrate that SLT-VEGF/L and SLT-VEGF/S molecules can enter cells via KDR/flk-1 receptors, and SLT/L or SLT/S moieties of said molecules can effectively cause cytotoxic and/or cytostatic effects in growing endothelial cells overexpressing KDR/flk-1 receptors, but not in endothelial cells that express low numbers of KDR/flk-1

20 receptors or quiescent endothelial cells.

These results demonstrate the possibility of using SLT-VEGF/L and SLT-VEGF/S proteins to target selectively growing endothelial at the sites of angiogenesis that are known to overexpress KDR/flk-1 receptors, without affecting normal endothelial cells or other types of cells that express either low number or none receptors for VEGF proteins,

25 thereby minimizing undesired side effects that might arise from interaction with not-

targeted cells. Therefore, SLT-VEGF/L and SLT-VEGF/S proteins are given a narrow spectrum of cellular interactions via specific binding to surface VEGF receptors in cells overexpressing said receptors thereby targeting SLT/L and SLT/S to primarily growing endothelial cells at the sites of angiogenesis.

5 Furthermore, using SLT-VEGF/L and SLT-VEGF/S constructs we have demonstrated that:

(i) SLT-VEGF/L and SLT-VEGF/S but not SLT-VEGF/Lci proteins retain the ability to inhibit protein synthesis.

10 (ii) SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S proteins bind to cellular KDR/flk-1 receptors and induce tyrosine autophosphorylation of said receptors.

(iii) SLT-VEGF/L protein is cytotoxic protein inducing death of growing endothelial cells that overexpress KDR/flk-1 receptors, but not endothelial cells that express low number of KDR/flk-1 receptors, or quiescent endothelial cells, while SLT-VEGF/S is mostly cytostatic protein causing growth inhibition.

15 The compositions for use in inhibition of endothelial cell growth in order to inhibit angiogenesis comprise a fusion protein, in combination with a pharmaceutically acceptable diluent or carrier. The compositions according to the invention will in practice normally be administered by intravenous injection, continuous infusion, although other methods, such as parenteral injection or intramuscular injection may also be used.

20 Compositions for injection can be provided in unit dose form and can take a form such as solution and can contain formulating agents, such as stabilizing agents, buffers, and the like.

The invention is further described by the following Examples, but is not intended to be limited by the Examples. All parts and percentages are by weight and all 25 temperatures are in degrees Celsius unless explicitly stated otherwise.

EXAMPLES

EXAMPLE 1 - CONSTRUCTION OF DNA SEQUENCES ENCODING SLT-VEGF/L
and SLT-VEGF/S FUSION PROTEINS

5

GENERAL DESCRIPTIONS

Bacterial Strains, Plasmids, and Mammalian Cells

E. coli strain DH5 α is commercially available from Life Technologies, Inc. (USA)

E. coli strains BL21(DE3)pLysS and Origami(DE3)pLysS are commercially available

10 from Novagen. Vector pET32(a) for bacterial expression of recombinant proteins with a terminal extension containing His-tag, S-tag, and thioredoxin is commercially available from Novagen (USA). Plasmid pLen-121 containing the DNA sequence encoding the 121-residue form of human VEGF has been described in US Patent No. 5,219,739, herein incorporated by reference in its entirety, and was obtained from Dr. J. Abraham (Scios

15 Nova, Inc., USA). Plasmid pJB144 containing a sequence for VT1/SLT holotoxin was obtained from Dr. J. Brunton (Samuel Lunenfeld Research Institute, Toronto, Canada).

Plasmids pBalPst (empty vector) and pBalPst/KDR encoding KDR/flk-1 receptor were obtained from Dr. B. Terman (Albert Einstein School of Medicine, New York City, USA).

Porcine aortic endothelial (PAE) cells and 293 human primary embryonic kidney cells

20 (293) were obtained from American Type Culture Collection (USA). PAE cells expressing 2-3x10⁵ KDR/flk-1 per cell (PAE/KDR), PAE cells transfected with pBalPst plasmid (PAE/V), human umbilical vein endothelial (HUVE) cells and MS1 mouse endothelial cells were obtained from Dr. B. Terman (Albert Einstein School of Medicine, New York City, USA). 293 cells overexpressing VEGFR-2 (293/KDR) and PAE cells

25 expressing a low number of KDR/flk-1 (PAE/KDR_{low}) were constructed by transfection of the corresponding parental cells with pBalPst/KDR plasmid using TransIT-LT1 reagent

(PanVera Corporation, USA), followed by selection in the presence of 0.375 g/ ml puromycin. A clone of 293/KDR cells chosen for this study expressed 2.5×10^6 VEGFR-2 per cell according to Scatchard's analysis of ^{125}I -VEGF165 binding. Expression levels of VEGFR-2 in PAE/KDR and PAE/KDR_{low} cells were estimated by Western blot analysis with 293/KDR cells serving as a standard. The immunoblots were probed with rabbit polyclonal anti-VEGFR-2 serum obtained from Dr. B. Terman (Albert Einstein School of Medicine, New York City, USA). PAE, 293 cells, and their derivatives were maintained in DMEM supplemented with 10% fetal bovine serum (Gemini, Inc., USA), 2 mM L-glutamine and antibiotics. Low passage number HUVE cells (3rd-7th passages) were grown in gelatin-coated flasks in DMEM with 20% FBS, 50 ng/ml basic fibroblast growth factor, 100 µg/ml heparin, 2 mM L-glutamine and antibiotics. MS1 cells are grown in DMEM with 5% FBS, 4 mM L-glutamine and antibiotics. All cell lines were cultivated at 37°C, 5% CO₂.

15 DNA Manipulations

The restriction and modification enzymes employed herein are commercially available from the usual sources and were used according to manufacturer's instructions. The sequencing of the different DNA constructs was done at Macromolecular Resources (Department of Biochemistry and Molecular Biology, Ft. Collins, CO, USA). Competent cells, transformation, and bacterial media were prepared according to Sambrook et al. (J. Sambrook, E. F. Fritsch and T. Maniatis. (1989) Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY), or according to the manufacturer's instructions. Purification of plasmids was done using Wizard Plus SV Minipreps or Maxipreps DNA Purification Systems (Promega, USA) according to the manufacturer's instructions. Further purification of DNA as well as purification of DNA

from agarose gels was done using the Geneclean Spin kit (Bio 101, USA) according to the manufacturer's instructions.

5 SUB-CLONING OF 121-RESIDUE ISOFORM OF HUMAN VEGF INTO A PET32(A) EXPRESSION VECTOR.

Primers for amplification of DNA encoding 121-residue isoform of human VEGF

Primers for human VEGF DNA amplifications were synthesized by GeneLink
10 (USA). The primer corresponding to the "sense" strand (SEQ ID NO:1) included a Stu I restriction site immediately upstream of the DNA codon for amino acid-1 of the mature 121-residue isoform of VEGF. The primer corresponding to the "antisense" strand (SEQ ID NO:2) complemented the coding sequence of the DNA encoding the carboxyl end of the mature peptide, including a translation stop codon after the sequence encoding mature
15 VEGF, and introduced an Xho I restriction site downstream of the VEGF-encoding DNA and the stop codon.

20 5'- TAAGGCCTATGGCAGAAGGAGGAGGG -3' (SEQ ID NO:1)

5'- ACTCGAGTCACCGCCTCGGCTTGTCA -3' (SEQ ID NO:2)

PCR to Amplify DNA Encoding 121-residue isoform of human VEGF

The human VEGF cDNA was amplified by PCR from the pLen-121 plasmid containing the sequence for the 121-residue isoform of human VEGF. Ten nanograms of template DNA were mixed in a 0.1 ml reaction mixture, containing 10 pmol of each
25 oligonucleotide, 0.2 mM of each dNTP and 2 U of Pfu polymerase (Stratagene, USA) in Pfu buffer (Stratagene, USA). Incubations were done in a DNA GenAmp PCR System 2400 (Perkin Elmer Cetus, USA). One cycle included a denaturation step (94°C for 1

min.), an annealing step (65°C for 1 min), and an elongation step (72°C for 1 min). The amplified DNA was digested with Stu I and Xho I and purified with the Geneclean Spin kit (BIO 101, USA).

5 pET32-VEGF121 Plasmid Construction

Amplified DNA forms described above were ligated into a pET32(a) vector using Xho I site from multiple cloning site of the vector and Stu I site, which was constructed by treating the vector as follows: pET32(a) DNA was linearized with Nco I restriction endonuclease, and one of produced recessed terminus was partially filled with cytidine using DNA polymerase I Large (Klenow) fragment. The construct was purified with the Geneclean Spin kit (BIO 101, USA) and single-stranded overhangs were removed with mung bean nuclease. The resulted construct was digested with Xho I restriction endonuclease and purified with the Geneclean Spin kit (BIO 101, USA). The ligation was accomplished such that the first amino acid of the mature 121-residue isoform of VEGF became the first amino acid after 10 an enterokinase cleavage site provided by vector. The resulting plasmid was designated pET32-txVEGF121 and was transformed into DH5 α competent cells (Life Technologies, USA) according to the manufacturer's instructions. The bacterial culture containing the desired plasmid was grown further in order to obtain large preparations of isolated plasmid using methods described above.

20 The thioredoxin (tx) gene was removed from the pET32-txVEGF121 by digestion of the purified plasmid DNA with restriction endonuclease Nde I, followed by intramolecular ligation of the linearized plasmid DNAs with T4 ligase. The resulting plasmid was designated pET32-VEGF121 and was transformed into DH5 α competent cells (Life Technologies, USA) according to the manufacturer's instructions. The bacterial culture containing the 25 desired plasmid was grown further in order to obtain large preparations of isolated plasmid

using methods described above. Plasmid pET-VEGF121 DNA encodes a 36 amino-acids full-length N-terminus, containing His-tag (6 amino acids), trinbin cleavage site (6 amino acids), S-tag (15 amino acids), a 6-amino acid full-length connecting peptide containing enterokinase cleavage site, and 1 to 121 amino acids of the mature VEGF121 protein (Fig. 5 1).

Sub-Cloning of Full-Length, Truncated and Mutant Forms of SLT Subunit A into pET32-VEGF121 Vector

10 1. Primers for amplification of DNA encoding full-length (L) and truncated (S) forms of SLT subunit A.

Primers for DNA amplifications of DNA encoding L and S forms of SLT subunit A were synthesized by GeneLink (USA). The primers corresponding to the "sense" 15 strands of full-length (293-residue) SLT form (SEQ ID NO:3) and truncated (202-residue) SLT form (SEQ ID NO:4) included Bgl II restriction sites upstream of the DNA codon for amino acid-1 and amino acid-62, respectively, of the mature SLT subunit A. In order to clone SLT molecules in frame with the first Met of VEGF121 an additional G was inserted between the Bgl II sites and the ORFs of SLT:

20

5'- CCGAGATCTGAAGGAATTACCTTAGAC -3' (SEQ ID NO:3)

5'- CCCAGATCTGCTACGGCTTATTGTTGAACG -3' (SEQ ID NO:4)

The primer corresponding to the "antisense" strand of full-length SLT form 25 complemented the coding sequence of the SLT DNA encoding the carboxyl end of the mature SLT subunit A right upstream the stop codon (SEQ ID NO:5). The primer corresponding to the "antisense" strand of truncated SLT form complemented the coding sequence of the DNA encoding the DNA codons for amino acids 258-264 of SLT subunit

A (SEQ ID NO:6). Both primers introduced Kpn I restriction sites downstream of the SLT-encoding DNA:

5'- ATAGGTACCACTGCTAATAGTTCTGCG -3' (SEQ ID NO:5)

5 5'- ATAGGTACCATCTGCCGGACACATAGAAG -3' (SEQ ID NO:6)

PCR to Amplify Full-length and Truncated Forms of SLT Subunit A

DNA encoding full-length and truncated forms of SLT Subunit A were amplified by PCR from the plasmid pJB144 containing VT1/SLT holotoxin. Ten nanograms of template DNA were mixed in a 0.1 ml reaction mixture, containing 10 pmol of each oligonucleotide, 0.2 mM dNTPs and 2 U of Vent DNA polymerase (New England Biolabs, USA) in Vent buffer. Incubations were done in a DNA GenAmp PCR System 2400 (Perkin Elmer Cetus, USA). One cycle included a denaturation step (94°C for 30 sec.), an annealing step (58°C for 1 min), and an elongation step (72°C for 1 min 20 sec.).

15 After 25 cycles, a 10 µl aliquot of each reaction was run on a 1% agarose gel to verify the correct size of the amplified product. The amplified DNA forms were digested with Bgl II and Kpn I restrictionases and purified with the GeneClean Spin kit (BIO 101, USA).

pET32-VEGF121-SLT/L and pET32-VEGF121-SLT/S Plasmids Construction

20 Amplified SLT DNA forms described above were ligated into pET32-VEGF121 vector that had been treated with Bgl II and Kpn I restrictionases and purified as described above. The resulting plasmids containing DNA encoding full-length and truncated SLT forms were designated pET32-VEGF121-SLT/L and pET32-VEGF121-SLT/S, respectively, and transformed into DH5 α competent cells (Gibco, USA) according to the manufacturer's instructions. The clones were screened, purified, characterized and

propagated as described above. The DNA fragments in pET32-VEGF121-SLT/L and pET32-VEGF121-SLT/S plasmids were sequenced from T7 promoter to nucleotide 203 in the coding sequences of SLT in order to confirm that they contain expected SLT DNA sequences.

5

Plasmid pET-VEGF121-SLT/L DNA encodes a 36 amino-acids full-length N-terminus, containing His-tag (6 amino acids), trinobin cleavage site (6 amino acids), S-tag (15 amino acids), the entire SLT subunit A (293 amino acids), a 6-amino acid full-length connecting peptide containing enterokinase cleavage site, and 1 to 121 amino acids of the mature VEGF121 protein (Fig. 1). Plasmid pET-VEGF121-SLT/S is identical to the plasmid pET-VEGF121-SLT/L, but instead the DNA encoding the entire SLT subunit A it contains the DNA encoding a 202-amino acid fragment of this subunit from amino acid 62 to amino acid 264 (Fig. 1).

15 Construction of plasmid for expression of catalytically inactive SLT-VEGF/Lci protein.

Site-specific mutagenesis of SLT-1 A-subunit encoded by pET32/SLT-VEGF/L was done using GeneEditor™ *in vitro* Site-Directed Mutagenesis System (Promega). Two mutagenic primers were designed to introduce three point mutations (underlined): Y114S (SEQ ID NO:7), and E167Q and R170L (SEQ ID NO:8):

5'-ACGTGGTAGAGCTACTGTCACC-3' (SEQ ID NO:7)

5'-TTGCCGAAAAAGTAAGCTTGAGCTGTCACAG-3' (SEQ ID NO:8)

25 The Y114S and R170L mutations were confirmed by sequencing of mutated DNA isolated from two clones. The E167Q mutation was not detected in either clone. The

resulting plasmid designated pET32-VEGF121-SLT/Lci is identical to plasmid pET-VEGF121-SLT/L, but instead the DNA encoding wild type SLT subunit A it contains the DNA encoding a double mutant (Y114S and R170L) of this subunit (Fig. 1).

EXAMPLE 2 - EXPRESSION AND PURIFICATION OF RECOMBINANT SLT-
5 VEGF/L, SLT-VEGF/Lci, AND SLT-VEGF/S FUSION PROTEINS

A. Expression of SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S proteins

10 1. Expression of SLT-VEGF/L and SLT-VEGF/S proteins in E. coli BL21(DE3)pLysS

The pET-VEGF121-SLT/L and pET-VEGF121-SLT/S transformed E. coli cells BL21(DE3)pLysS (Novagen, USA) were grown under conditions in which the expression of the fusion proteins is repressed by the lac repressor to an O.D. in or at the middle of the 15 log phase of growth after which IPTG (isopropyl-D-thiogalactoside, Life Technologies, USA) was added to induce expression of the fusion protein-encoding DNA.

To generate a large-batch culture of pET-VEGF121-SLT/L and pET-VEGF121-SLT/S transformed E. coli cells, an overnight cultures (lasting approximately 16 hours) of BL21(DE3)pLysS E. coli cells transformed with the plasmids pET-VEGF121-SLT/L and 20 pET-VEGF121-SLT/S, respectively, in LB broth containing 50 mg/L ampicillin and 34 mg/L chloramphenicol was diluted 1:100 into a flask containing 100 ml LB broth with 50 mg/L ampicillin and 34 mg/L chloramphenicol. Cells were grown with shaking at 37°C until the optical density at 600 nm reached 0.5 as measured in a spectrophotometer (Ultrospec 1000, Pharmacia Biotech, USA).

25 In the second step, fusion protein expression was induced by the addition of IPTG (Life Technologies, USA) to a final concentration of 1 mM. Induced cultures were grown for 3.5 additional hours for SLT-VEGF/L and 2 additional hours for SLT-VEGF/S; then harvested by centrifugation (25 min., 5000xg). The cell pellets were resuspended in the

ice-cold buffer A (50 mM Tris-HCl, pH 7.5, 0.1 M MgCl₂, 1% Nonidet P 40, 0.1 M DTT, 200 mg/L PMSF, 25 mg/L antitrypsin, 50 mg/L leupeptin, 25 mg/L aprotinin). After five cycles of freezing and thawing, DNase was added to each of the cell suspensions to 50 U per ml. The suspensions were incubated for 20 min at room temperature, then centrifuged 5 at 5,000xg for 30 min at 4°C. Analysis of distribution of fusion proteins designated as SLT-VEGF/L and SLT-VEGF/S between soluble fractions and inclusion bodies of IPTG-induced bacteria indicated that SLT-VEGF/L and SLT-VEGF/S fusion proteins were present in the inclusion bodies (Fig.2, panel A).

The inclusion body pellets were washed with the buffer containing 20 mM Tris-10 HCL, pH 8.0, 0.5 M NaCl, 5 mM imidazole, and solubilized in 8 M urea, followed by sonication of the solutions for 5-10 min in an ice-cold water sonicator (FC 14, Fisher Sci., USA). The protein solutions were clarified by centrifugation at 14xg for 10 min at 4°C and the supernatants were collected and dialyzed against a 1000-fold volume of the buffer containing 10 mM Tris-HCl, pH 8.0, 150 mM NaCl, 0.05% Nonidet P 40 for 16 hours at 15 4°C. The SLT-VEGF/L and SLT-VEGF/S fusion protein solutions were supplemented with 10% glycerol and stored in aliquots at -20°C. The concentrations of recombinant proteins were determined with S-tag assay kit (Novagen, USA) according to manufacturer's instructions.

2. Expression of SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S proteins in E. coli Origami(DE3)pLysS

The pET-VEGF121-SLT/L, SLT-VEGF/Lci, and pET-VEGF121-SLT/S transformed E. coli cells Origami(DE3)pLysS (Novagen, USA) were grown under conditions in which the expression of the fusion proteins is repressed by the lac repressor to an O.D. in or at the middle of the log phase of growth after which IPTG (Life Technologies, USA) was added to induce expression of the fusion protein-encoding DNA.

To generate a large-batch culture of pET-VEGF121-SLT/L, pET-VEGF121-SLT/L/ci, and pET-VEGF121-SLT/S transformed E. coli cells, an overnight cultures (lasting approximately 16 hours) of Origami(DE3)pLysS E. coli cells transformed with the plasmids pET-VEGF121-SLT/L, or pET-VEGF121-SLT/L/ci, or pET-VEGF121-SLT/S, respectively, in LB broth containing 50 mg/L ampicillin, 34 mg/L chloramphenicol, 12.5 mg/L tetracycline, and 15 mg/L kanamycin was diluted 1:100 into a flask containing 100 ml LB broth with 50 mg/L ampicillin, 34 mg/L chloramphenicol, 12.5 mg/L tetracycline, and 15 mg/L kanamycin. Cells were grown with shaking at 37°C until the optical density at 600 nm reached 0.4 measured in a spectrophotometer (Ultrospec 1000, Pharmacia Biotech, USA).

In the second step, fusion protein expression was induced by the addition of IPTG (Life Technologies, USA) to a final concentration of 1 mM. Induced cultures were grown for 4 additional hours at 30°C; then harvested by centrifugation (25 min, 5000xg). The cell pellets were resuspended in the ice-cold buffer A (50 mM Tris-HCl, pH 7.5, 0.1 M MgCl₂, 1% Nonidet P 40, 0.1 M DTT, 200 mg/L PMSF, 25 mg/L antitrypsin, 50 mg/L leupeptin, 25 mg/L aprotinin). After five cycles of freezing and thawing, DNase was added to each of the cell suspensions to 50 U per ml. The suspensions were incubated for 20 min at room temperature; then centrifuged at 5,000xg for 30 min at 4°C. Analysis of

distribution of fusion proteins designated as SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S between soluble fractions and inclusion bodies of IPTG-induced bacteria indicated that SLT-VEGF/L and SLT-VEGF/S these proteins were present in the inclusion bodies (Fig. 2, panel B for SLT-VEGF/L and SLT-VEGF/S).

5 The inclusion body pellets were washed with the buffer containing 20 mM Tris-HCl, pH 8.0, 0.5 M NaCl, 5 mM imidazole, and solubilized in 8 M urea, followed by sonication of the ice-cold solutions for 20-30 sec in a sonicator VirSonic 475(VirTis, USA)operated at 40-50% of output power. The protein solutions were clarified by centrifugation at 14,000xg for 10 min at 4°C and the supernatants were collected and
10 dialyzed against a 1000-fold volume of the buffer containing 10 mM Tris-HCl, pH 8.0, 150 mM NaCl, 0.01% Brij-35 for 2 hours at 4°C and then for 16 hours at 4°C against a fresh 1000-fold volume of the same buffer. SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S fusion protein obtained through this procedure were characterized by SDS-PAGE (Fig. 2, panel C). The SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S fusion protein
15 solutions were supplemented with 5% glycerol and stored in aliquots at -70°C. The concentrations of recombinant proteins were determined with S-tag assay kit (Novagen, USA) according to manufacturer's instructions.

B. Expression of VEGF121 in E. coli BL21(DE3)pLysS

20 The pET32-VEGF121 transformed E. coli cells BL21(DE3)pLysS were grown and expression of recombinant VEGF designated as VEGF121 was induced under conditions described above. Recombinant VEGF121 was recovered from inclusion bodies as described above and supernatant was dialyzed against a 1000-fold volume of the buffer

containing 10 mM Tris-HCl, pH 8.0, 150 mM NaCl for 16 hours at 4°C. The VEGF121 protein solutions were supplemented with 10% glycerol and stored in aliquots at -20°C. The concentrations of recombinant VEGF121 proteins were determined with S-tag assay kit (Novagen, USA) according to manufacturer's instructions.

5

EXAMPLE 3 - BIOCHEMICAL ACTIVITIES OF SLT-VEGF/L AND SLT-VEGF/S FUSION PROTEINS

10 A. Inhibitory Effect of SLT-VEGF/L and SLT-VEGF/S Fusion Proteins on Cell-Free Protein Synthesis

Abilities of SLT-VEGF/L and SLT-VEGF/S recombinant fusion proteins, obtained above, to inhibit protein synthesis were tested in an in vitro assay measuring cell-free protein synthesis in a nuclease-treated rabbit reticulocyte lysate (Promega, USA). 5 µl of SLT-
15 VEGF/L, SLT-VEGF/S, VEGF121 solutions or storage buffer containing 10 mM Tris-HCL, 150 mM NaCl, 8 mM urea, 10% glycerol were added on ice to 20µl reaction mixtures that included 18 µl of rabbit reticulocyte lysate, 1 mM complete amino acid mix, 90 mM KCl, and firefly luciferase mRNA, 0.5 µg per assay. The final concentrations of SLT-VEGF/L, SLT-VEGF/S were 100 nM and a final concentration of VEGF121 was
20 1000 nM. The final concentrations of SLT-VEGF/L, SLT-VEGF/Lci, SLT-VEGF/S in dose-dependence experiments were as indicated in Figure 3, panel B. After a 90-min incubation at 30°C the reaction mixtures were diluted 20-fold with water containing 1 mg/ml BSA, and 2 µl of each mixture was assayed for firefly luciferase activity in Luciferase Reaction buffer (Promega, USA) using LumiOne luminometer (Bioscan, Inc,
25 USA) according to the manufacturer's instructions. Luciferase activity of the control reaction containing 100 nM VEGF121 was taken as 100%.

SLT-VEGF/L and SLT-VEGF/S fusion proteins at concentration 100 nM inhibited protein synthesis to the levels of 0.01% and 1.2% of control (Fig. 3, panel A). For comparison, recombinant VEGF121 protein at concentration as high as 1000 nM inhibited protein synthesis to the level of 48% of control (Fig. 3, panel A). Dose-dependence experiments 5 revealed that SLT-VEGF/L and SLT-VEGF/S inhibited protein synthesis by 90% at concentrations 0.04 nM and 2 nM, respectively, while SLT-VEGF/Lci did not inhibit protein synthesis (Fig. 3, panel C).

10 B. Induction of the Tyrosine Phosphorylation of Cellular KDR/flk-1 Receptors by SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S Fusion Proteins

Abilities of SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S recombinant fusion proteins obtained above to induce tyrosine phosphorylation of cellular KDR/flk-1 receptors, was tested using 293 cells stably transfected with KDR/flk-1 designated as 15 293/KDR cells. About 50,000 293/KDR cells per well were plated onto 24-well plates in 1 ml DMEM (Life Technologies, USA) supplemented with 10% fetal calf serum (Gemini, USA) and incubated overnight at 37°C, 5% CO₂. Next day, the cells were washed once with phosphate buffered saline and shifted to DMEM serum-free for 4 hours at 37°C. Subsequently, the medium was changed to serum-free DMEM supplemented with 0.1 mM 20 sodium orthovanadate, 100 ng/ml bovine serum albumine, 25 mM HEPES pH 7.2, and the cells were incubated for 20 min at 37°C followed by a 20-min incubation at 4°C. Then the cells were incubated with SLT-VEGF/L, or SLT-VEGF/Lci, or SLT-VEGF/S, or VEGF121 for 1 hour at 4°C followed by 8 min at 37°C. Then the cells were rinsed once with ice-cold phosphate buffered saline containing 0.1 mM sodium orthovanadate, 25 solubilized in sample buffer containing 0.05 M Tris-HCl, pH 6.8, 2.5% SDS, 7.5%

glycerol, 5 mM EDTA, 50 mM DTT, 0.025% Bromophenol Blue, and analyzed by Western blotting. Cellular proteins were fractionated by SDS-PAGE on 7.5% gels and were transferred to nitrocellulose (BioRad, USA) using a semi-dry system 2117 Multiphor II (LKB, Sweden), as described by the manufacturer. Western blots were processed and 5 probed with anti-phosphotyrosine RC20:HRP conjugate (Transduction Lab, USA) at dilution 1:2,000 according to the manufacturer's instructions. A chemiluminescence-based system (ECL, Amersham, USA) was used for bands detection. SLT-VEGF/L, SLT-VEGF/Lci, and SLT-VEGF/S proteins induced KDR/flk-1 tyrosine phosphorylation in a dose-dependent manner at the same concentration range as VEGF121. (Fig.4).

10

EXAMPLE 4 - CYTOTOXIC AND CYTOSTATIC EFFECTS OF SLT-VEGF/L AND SLT-VEGF/S FUSION PROTEINS

15 A. Effects of VEGF-SLT/L and VEGF-SLT/S Fusion Proteins on Growing Endothelial Cells Overexpressing KDR/flk-1 Receptors and Growing Endothelial Cells That Do Not Express KDR/flk-1 Receptors

Abilities of VEGF-SLT/L and VEGF-SLT/S fusion proteins obtained above to affect growing endothelial cell were tested using porcine aortic endothelial cells stably 20 transfected with KDR/flk-1 (designated as PAE/KDR cells) and porcine aortic endothelial cells stably transfected with the empty vector (designated as PAE/V cells). About 5,000 cells per well were plated onto 24-well plates in 1 ml DMEM (Life Technologies, USA) supplemented with 10% fetal calf serum (Gemini, USA) and incubated overnight at 37°C, 5% CO₂. Next day the media was changed to fresh DMEM supplemented with 10% fetal 25 calf serum, and SLT-VEGF/L or SLT-VEGF/S at final concentrations from 0.078 nM to 2.5 nM. Plates were incubated for 72 hours, at 37°C, 5% CO₂. Following the incubation period, wells were washed with phosphate buffered saline, cells were detached with

trypsin solution (Life Technologies, USA) and counted in a Coulter Counter (Coulter Corporation, USA) according to the manufacturer's instructions. SLT-VEGF/L and SLT-VEGF/S fusion proteins inhibited growth of PAE/KDR cells in a dose-dependent manner with IC₅₀~0.15 nM while growth of PAE/V that do not express KDR/flk-1 receptors was 5 not affected (Fig. 5). At the low nanomolar concentration range SLT-VEGF/L proteins were cytotoxic, killing virtually all PAE/KDR cells after overnight exposure to concentration as low as 2.5 nM. In contrast, SLT-VEGF/S proteins at the low nanomolar concentrations remained mostly cytostatic with few dead cells observed at concentration 2.5 nM.

10 Catalytically inactive SLT-VEGF/Lci fusion protein did not affect growth of PAE/KDR and PAE/V cells (Fig. 5, Panel C) indicating that cytotoxic and cytostatic effects of SLT-VEGF/L and SLT-VEGF/S fusion proteins are due to the ribosome inactivating activity of SLT moiety.

Cytotoxic activity of SLT-VEGF/L manifested itself in rapid induction of 15 apoptosis in PAE/KDR cells as judged by such hallmarks of apoptosis as DNA fragmentation and cleavage of -fodrin (Fig. 6) To detect apoptotic DNA fragmentation PAE/KDR cells were plated onto 6-well plates at a density of 2x10⁵ cells/well and exposed to 5 nM SLT-VEGF/L 24 hr later. After indicated periods of time DNA was isolated from cell lysates and fractionated on 1.5% agarose gel. To detect cleavage of 20 kDa -fodrin into 150 kDa and 120 kDa fragments, PAE/KDR cells were plated onto 24-well plates at a density of 4x10⁴ cells/well and exposed to 2.5 nM SLT-VEGF/L 20 hr later for indicated periods of time. -fodrin and its fragments were detected by Western blot analysis of the lysates of treated cells, using anti- -fodrin antibody (Chemicon, USA).

B. Effects of VEGF-SLT/L Fusion Protein on Endothelial Cells Expressing Low Number of KDR/flk-1 Receptors and on Quiescent Endothelial Cells

Endothelial cells in the normal vasculature express low number of KDR/flk-1 receptors. To minimize potential negative side effects a useful toxin-VEGF fusion protein should not be toxic for endothelial cells with low number of KDR/flk-1 receptors or to quiescent endothelial cells that constitute normal vasculature. The ability of VEGF-SLT/L fusion protein obtained above to affect endothelial cells expressing low number of KDR/flk-1 receptors was tested with human umbilical vein endothelial (HUVE) cells expressing 30,000-50,000 KDR/flk-1 receptors per cell, PAE/KDR_{low} porcine aortic endothelial cells expressing ~5,000 KDR/flk-1 receptors per cell, and MS1 mouse endothelial cells expressing ~20,000 VEGFR-2/cells. HUVE, PAE/KDR_{low}, and MS1 cells were plated onto 24-well plates at densities of 5-10x10³ cells/well and exposed to 2.5 nM SLT-VEGF/L 20 hr later. Plates were incubated for 72 hours, at 37°C, 5% CO₂.

Following the incubation period, wells were washed with phosphate buffered saline (PBS), cells were detached with trypsin solution (Life Technologies, USA) and counted in a Coulter Counter (Coulter Corporation, USA) according to the manufacturer's instructions. SLT-VEGF/L fusion protein did not affect growing human HUVE, mouse MS1, and porcine PAE/KDR_{low} endothelial cells indicating that only a high level of KDR/flk-1 receptor expression may confer sensitivity to SLT-VEGF/L fusion protein on endothelial cells (Fig. 7, panel A).

The ability of VEGF-SLT/L fusion protein obtained above to affect quiescent endothelial cells was tested with quiescent PAE/KDR cells. About 5,000 PAE/KDR cells per well were plated onto 24-well plates in 1 ml DMEM (Life Technologies, USA) supplemented with 10% fetal calf serum (Gemini, USA) and incubated at 37°C, 5% CO₂. Cells reached confluence after approximately one week and were maintained as quiescent

for additional three days. After that the media was changed to fresh DMEM supplemented with 10% fetal calf serum, and SLT-VEGF/L at a final concentration of 20 nM. Plates were incubated for 72 hours, at 37°C, 5% CO₂. For comparison, growing PAE/KDR were exposed to 20 nM SLT-VEGF/L for 5 min; then shifted to fresh culture medium and

5 counted after 72 hrs. SLT-VEGF/L proteins did not affect quiescent PAE/KDR endothelial cells even after 72-hour exposure, while dramatically affected growing PAE/KDR even after a 5-min exposure.

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WHAT IS CLAIMED IS:

1. An isolated nucleic acid encoding a fusion protein comprising:
 - (1) the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version

5 thereof; and

- (2) human vascular endothelial growth factor, or a truncated or mutated version thereof;

wherein said fusion protein possesses ribosome inactivating activity.

10 2. The isolated nucleic acid of claim 1, wherein said fusion protein specifically binds to vascular endothelial growth factor receptors.

3. The isolated nucleic acid of claim 2, wherein said fusion protein is internalized by a cell which expresses said receptors.

15

4. The isolated nucleic acid of claim 3, wherein said internalization occurs by endocytosis.

5. An isolated polypeptide comprising:

20 (1) the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and

- (2) human vascular endothelial growth factor, or a truncated or mutated version thereof;

wherein said isolated polypeptide possesses ribosome inactivating activity.

25

6. The isolated polypeptide of claim 5, wherein said isolated polypeptide specifically binds vascular endothelial growth factor receptors.
7. The isolated polypeptide of claim 6, wherein said isolated polypeptide is internalized by a cell which expresses said receptors.
8. The isolated polypeptide of claim 5, wherein said internalization occurs by endocytosis.

10 9. An expression vector, comprising:

- (1) a nucleic acid encoding a fusion protein comprising the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and human vascular endothelial growth factor, or a truncated or mutated version thereof; and
- (2) a promoter sequence operably linked to said nucleic acid to allow expression of said nucleic acid.

10. The expression vector of claim 9, wherein said fusion protein specifically binds vascular endothelial growth factor receptors.

20 11. The expression vector of claim 10, wherein said fusion protein is internalized by a cell which expresses said receptors.

12. A bacterial cell transformed with the expression vector of claim 10.

13. A method of inactivating ribosomes in a cell, comprising the steps of:

(a) contacting a cell with a polypeptide comprising:

(1) the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and

5 (2) human vascular endothelial growth factor, or a truncated or mutated version thereof; under conditions which permit said polypeptide to be internalized into said cell and inactivate ribosomes in said cell.

14. The method of claim 13, wherein said fusion protein specifically binds vascular

10 endothelial growth factor receptors.

15. The method of claim 14, wherein said fusion protein is internalized by a cell which

expresses said receptors.

15 16. A composition for inhibiting endothelial cell growth in a patient, comprising:

(A) a fusion protein comprising the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and human vascular endothelial growth factor, or a truncated or mutated version thereof, said fusion protein possessing ribosome inactivating activity; and

20 (B) a pharmaceutically acceptable carrier.

17. A method of treating a patient suffering from a pathophysiological condition that depends on angiogenesis, comprising:

providing to said patient an effective amount of a composition comprising a fusion protein comprising the A subunit of Shiga-like bacterial toxin, or a truncated or mutated version thereof; and human vascular endothelial growth factor, or a truncated or mutated version thereof, said fusion protein possessing ribosome inactivating activity; and a pharmaceutically acceptable carrier.

5
18. The method of claim 17, wherein said method is used in combination with other 10 treatments for said pathophysiological condition.

19. The method of claim 17, wherein the efficacy of said treatment is enhanced by decrease in oxygen or nutrient supplies that would arise from damage to endothelium caused by said protein and pharmaceutical compositions.

15

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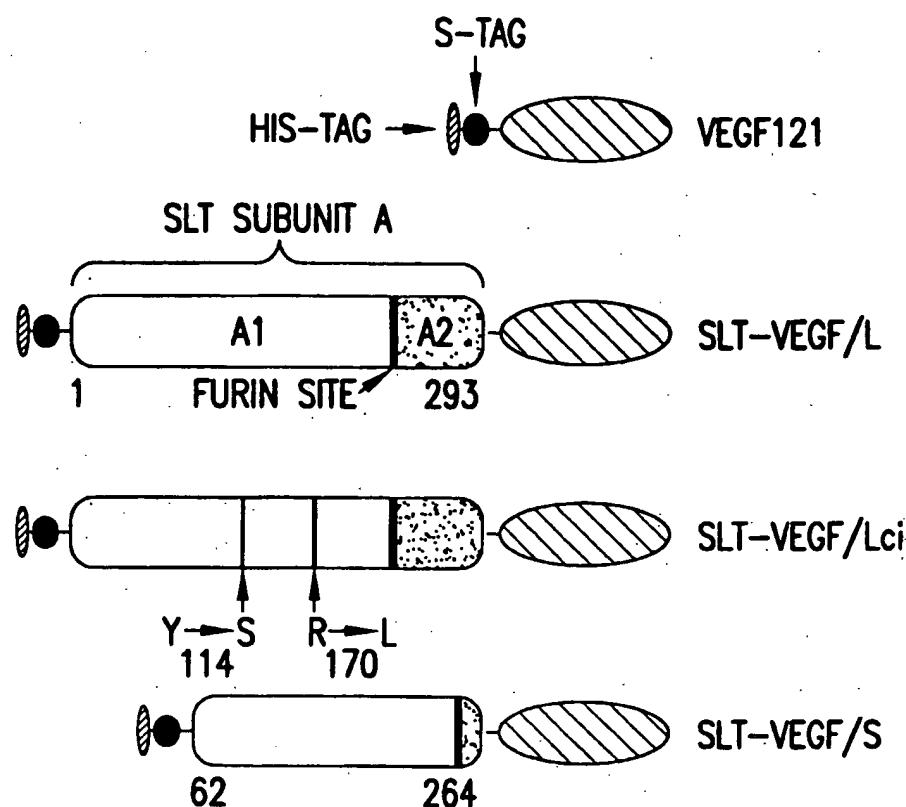


FIG.1

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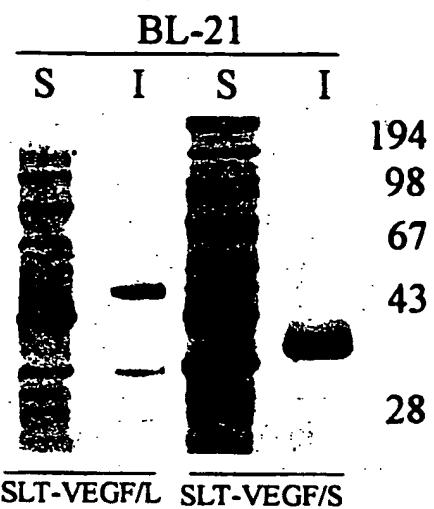


FIG.2A

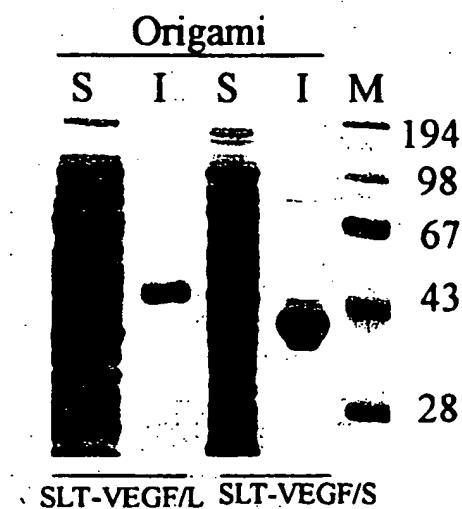


FIG.2B

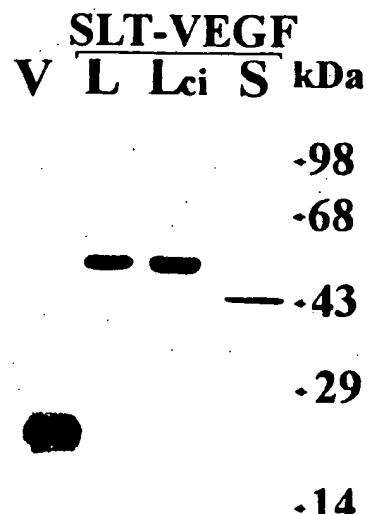


FIG.2C

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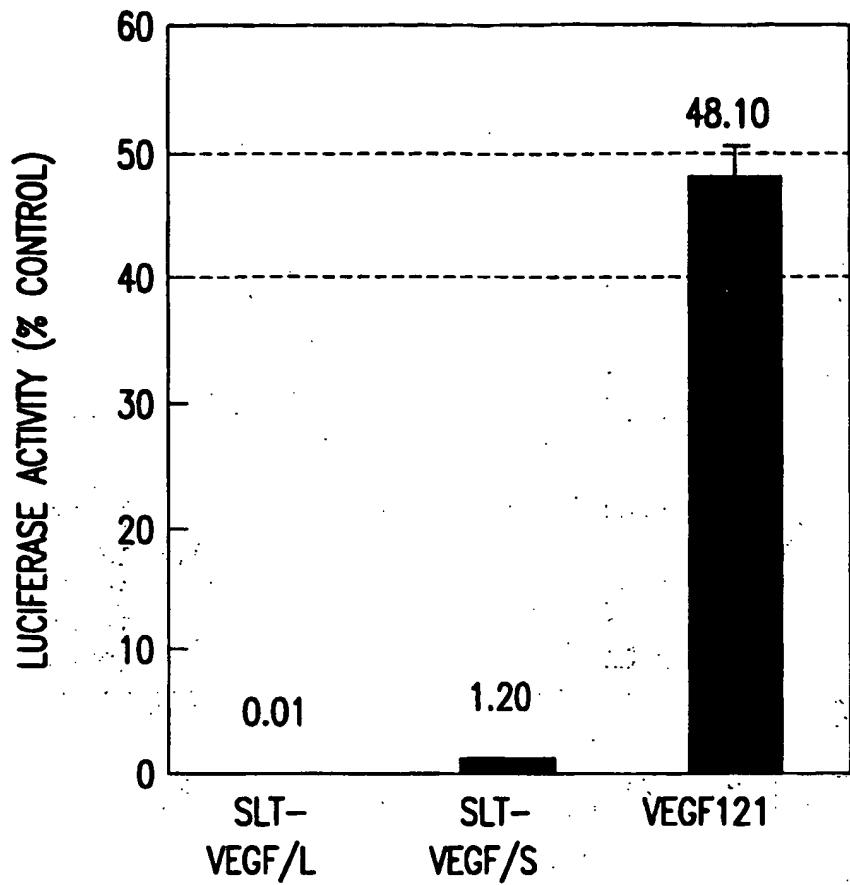


FIG.3A

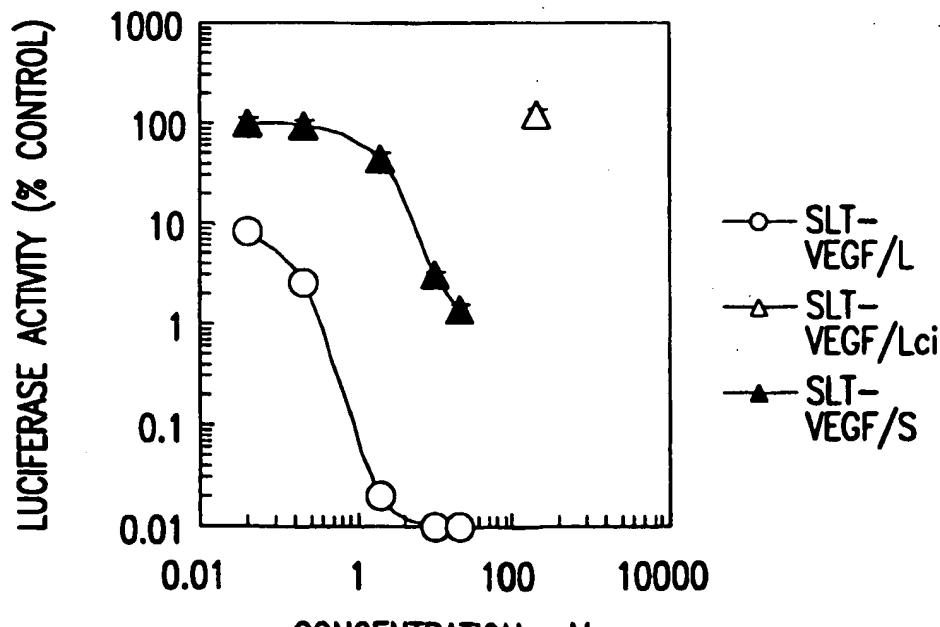


FIG.3B

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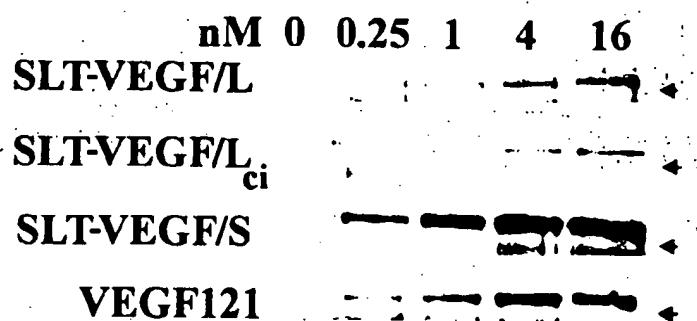


FIG.4

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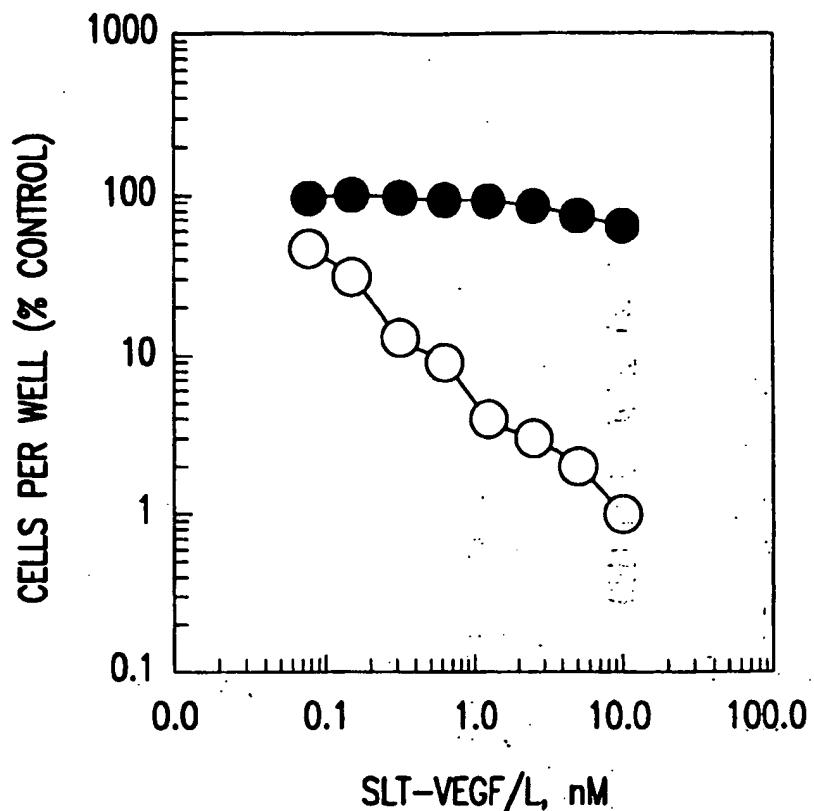


FIG.5A

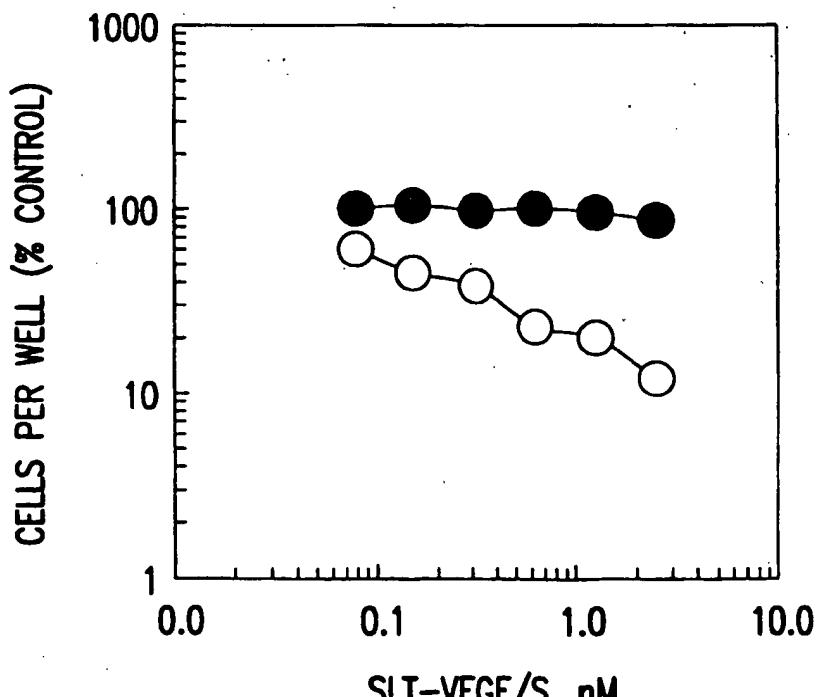


FIG.5B

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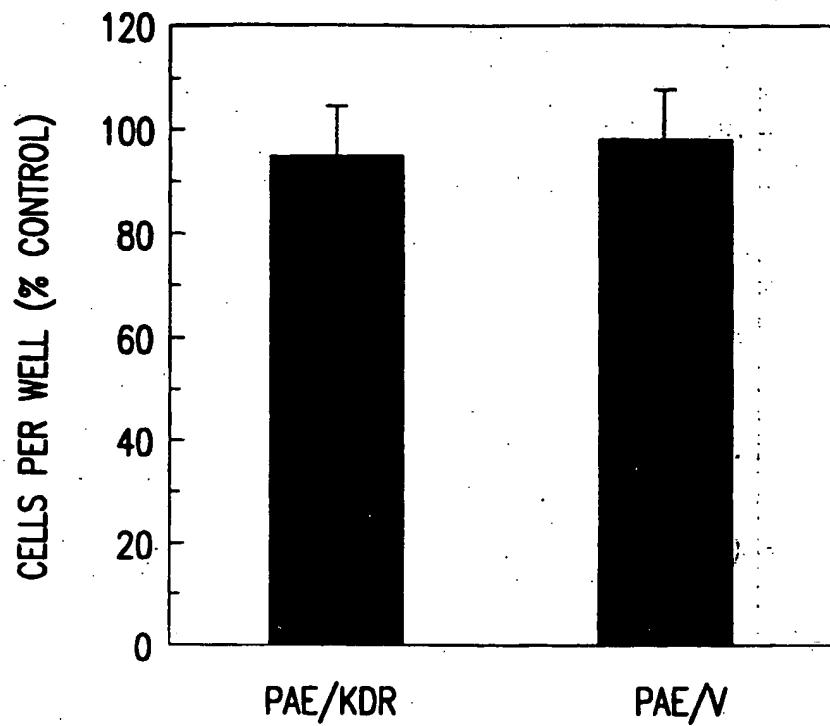


FIG.5C

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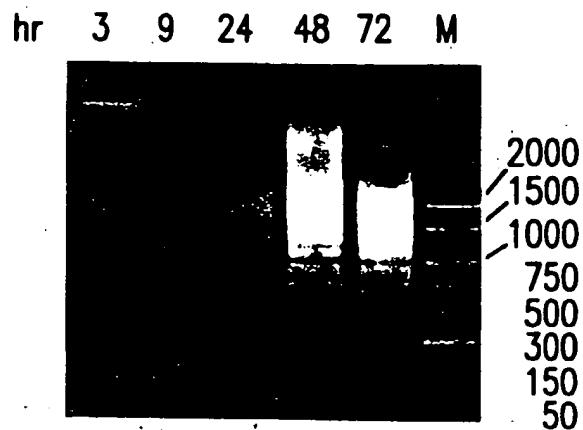


FIG. 6A

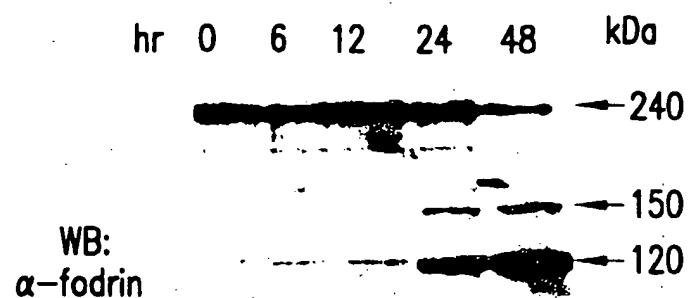


FIG. 6B

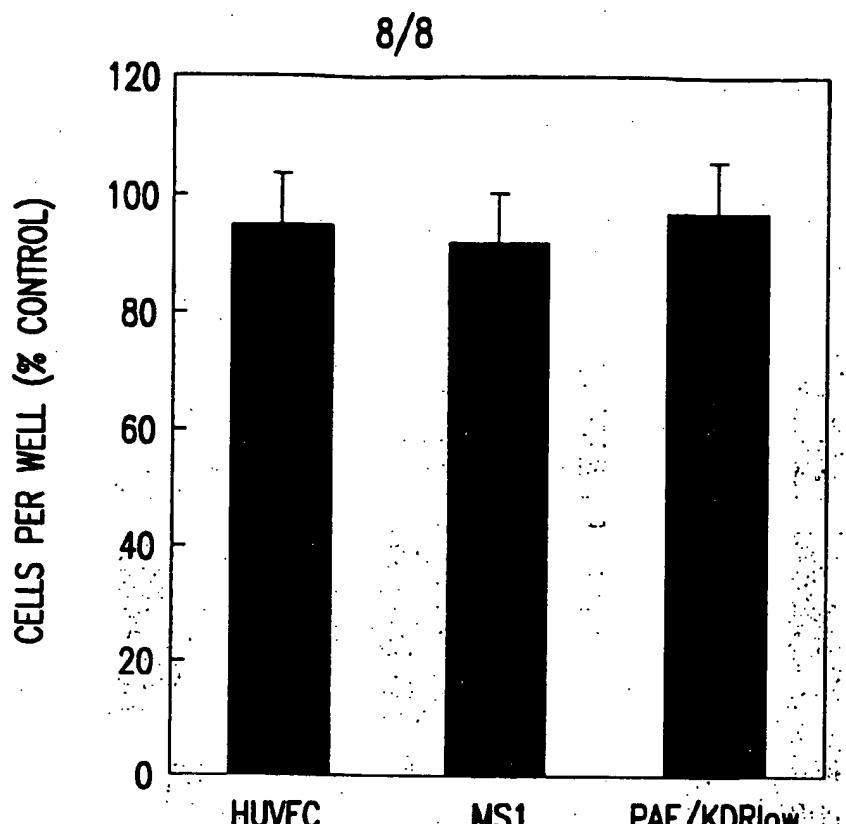


FIG. 7A

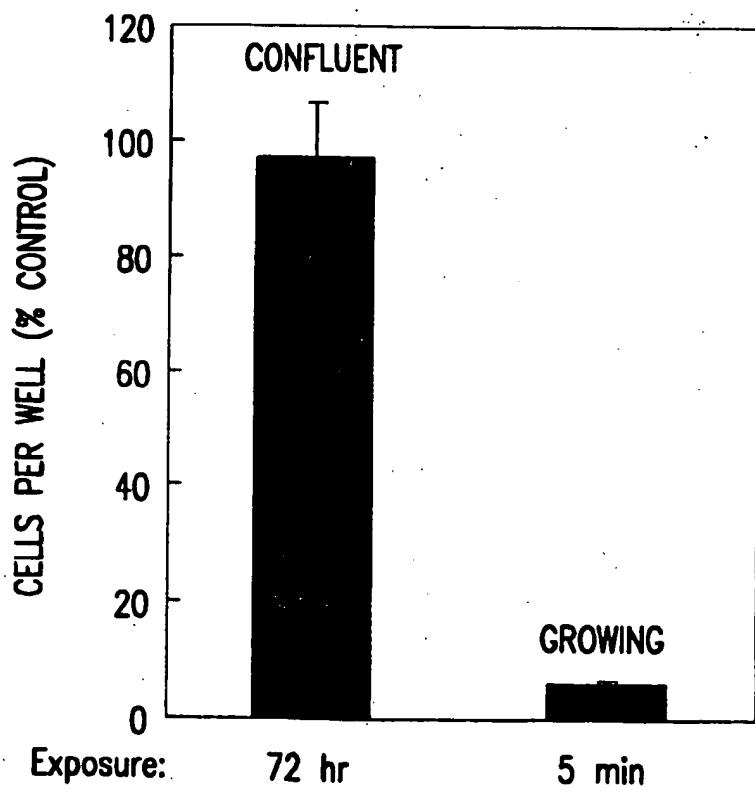


FIG. 7B

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Backer, Joseph M.

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32

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US01/08798

A. CLASSIFICATION OF SUBJECT MATTER

IPC(7) : C12N 15/00; C07K 14/00, 14/52, 14/245
 US CL : 435/69.1; 530/350, 351; 514/2

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 435/69.1; 530/350, 351; 514/2

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

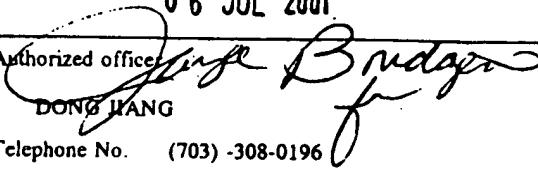
STN (medline, biosis, uspatfull, wpix)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	US 6,037,329 (BAIRD et al.) 14 March 2000, col. 2, 8, 9, 11, 20, 21, 47, 72.	1-18
Y	BRIGOTTI, M et al. The RNA-N-glycosidase activity of Shiga-like toxin I: Kinetic parameters of the native and activated toxin. Toxicon. September 1997, 35, No 9, pages 1431-1437, especially the first two sentences of the abstract, and the first two paragraphs.	1-18

Further documents are listed in the continuation of Box C. See patent family annex.

Special categories of cited documents:	"T"	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"A"	document defining the general state of the art which is not considered to be of particular relevance	
"E"	earlier document published on or after the international filing date	"X"
"L"	document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"Y"
"O"	document referring to an oral disclosure, use, exhibition or other means	
"P"	document published prior to the international filing date but later than the priority date claimed	"A"
		document member of the same patent family

Date of the actual completion of the international search	Date of mailing of the international search report
30 MAY 2001	06 JUL 2001
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231	Authorized officer  DONG JIANG
Faxsimile No. (703) 305-3230	Telephone No. (703) -308-0196

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US01/08798

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:

2. Claims Nos.: because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

Please See Extra Sheet.

1. As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

The additional search fees were accompanied by the applicant's protest.

No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US01/08798

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING

This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be searched, the appropriate additional search fees must be paid.

Group I, claim(s) 1-12, drawn to an isolated nucleic acid, an isolated polypeptide encoded thereby, and an expression vector and a bacterial cell containing such.

Group II, claim(s) 13-19, drawn to a method of inactivating ribosomes in a cell, a composition for inhibiting endothelial cell growth in a patient, and method of treating a patient using said composition.

The inventions listed as Groups do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons:

This authority considers that the main invention in the instant application comprises the first-recited product, an isolated nucleic acid encoding a fusion protein, and the first-recited method of using that product, namely a process of recombinant expression and purification of the encoded polypeptide. Also included in this group is the product made, namely the encoded polypeptide, an expression vector and a host cell containing the nucleic acid. The methods of inactivating ribosome, and treating a patient in group II do not share a special technical feature within the meaning of PCT Rule 13.2, and do not relate to a single invention concept within the meaning of PCT Rule 13.1.

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